



## Order form for MS Analysis

Name:	<input type="text"/>
Group:	<input type="text"/>
Institute:	<input type="text"/>
Street:	<input type="text"/>
Postal Code:	<input type="text"/>
Town:	<input type="text"/>
Email:	<input type="text"/>

**Send us a gelpicture where the bands or spots of interests are clearly marked**

How many samples should we analyse:

### From which organism is the sample:

- Human   
Yeast   
E. coli   
Drosophila   
Others

### Comments:

### Which method should we use:

- Protein identification by PMF (MALDI-TOF MS)  
 Protein identification by peptide sequencing (LC-ESI-MS/MS)  
 De Novo sequencing by ESI-MS/MS  
 Modifications  
 Mass determination of proteins/peptides

**Send the gel to:**

**ZMBH  
Mass Spectrometry  
Im Neuenheimer Feld 282  
69120 Heidelberg**

**For further questions call 06221-546895  
or Email: [MS-Service@zmbh.uni-heidelberg.de](mailto:MS-Service@zmbh.uni-heidelberg.de)**